

0590
10/2

#2 OIPE

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DATE: 11/01/2001

PATENT APPLICATION: US/09/964,238

TIME: 13:41:55

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 7 <130> FILE REFERENCE: 2676-4232US
 9 <140> CURRENT APPLICATION NUMBER: 09/964,238
 10 <141> CURRENT FILING DATE: 2001-09-26
 12 <150> PRIOR APPLICATION NUMBER: 09/449,285
 13 <151> PRIOR FILING DATE: 1999-11-24
 15 <150> PRIOR APPLICATION NUMBER: PCT/EP98/03193
 16 <151> PRIOR FILING DATE: 1998-05-28
 18 <150> PRIOR APPLICATION NUMBER: 97201645.5
 19 <151> PRIOR FILING DATE: 1997-06-02
 21 <160> NUMBER OF SEQ ID NOS: 27
 23 <170> SOFTWARE: PatentIn version 3.0
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 26 <211> LENGTH: 3006
 27 <212> TYPE: DNA
 28 <213> ORGANISM: Mus musculus
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148 35 40 45
150 Ser His Ser Gly Ser Tyr Ser Ser His Ile Ser Ser Lys Lys Cys Ile
151 50 55 60
153 Gly Leu Ile Ser Val Asn Gly Arg Met Arg Asn Asn Ile Lys Thr Gly
154 65 70 75 80
156 Ser Ser Pro Asn Ser Val Ser Ser Ser Pro Thr Asn Ser Ala Ile Thr
157 85 90 95
159 Gln Leu Arg Asn Lys Leu Glu Asn Gly Lys Pro Leu Ser Met Ser Glu
160 100 105 110
162 Gln Thr Gly Leu Leu Lys Ile Lys Thr Glu Pro Leu Asp Phe Asn Asp
163 115 120 125
165 Tyr Lys Val Leu Met Ala Thr His Gly Phe Ser Gly Ser Ser Pro Phe
166 130 135 140
168 Met Asn Gly Gly Leu Gly Ala Thr Ser Pro Leu Gly Val His Pro Ser
169 145 150 155 160
171 Ala Gln Ser Pro Met Gln His Leu Gly Val Gly Met Glu Ala Pro Leu
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180 Thr Glu Asp Ile Ser Lys Leu Lys Gly Tyr His Met Lys Asp Pro Cys
181      210      215      220
183 Ser Gln Pro Glu Glu Gln Gly Val Thr Ser Pro Asn Ile Pro Pro Val
184 225      230      235      240
186 Gly Leu Pro Val Val Ser His Asn Gly Ala Thr Lys Ser Ile Ile Asp
187      245      250      255
189 Tyr Thr Leu Glu Lys Val Asn Glu Ala Lys Ala Cys Leu Gln Ser Leu
190      260      265      270
192 Thr Thr Asp Ser Arg Arg Gln Ile Ser Asn Ile Lys Lys Glu Lys Leu
193      275      280      285
195 Arg Thr Leu Ile Asp Leu Val Thr Asp Asp Lys Met Ile Glu Asn His
196      290      295      300
198 Ser Ile Ser Thr Pro Phe Ser Cys Gln Phe Cys Lys Glu Ser Phe Pro
199 305      310      315      320
201 Gly Pro Ile Pro Leu His Gln His Glu Arg Tyr Leu Cys Lys Met Asn
202      325      330      335
204 Glu Glu Ile Lys Ala Val Leu Gln Pro His Glu Asn Ile Val Pro Asn
205      340      345      350
207 Lys Ala Gly Val Phe Val Asp Asn Lys Ala Leu Leu Leu Ser Ser Val
208      355      360      365
210 Leu Ser Glu Lys Gly Leu Thr Ser Pro Ile Asn Pro Tyr Lys Asp His
211      370      375      380
213 Met Ser Val Leu Lys Ala Tyr Tyr Ala Met Asn Met Glu Pro Asn Ser
214 385      390      395      400
216 Asp Glu Leu Leu Lys Ile Ser Ile Ala Val Gly Leu Pro Gln Glu Phe
217      405      410      415
219 Val Lys Glu Trp Phe Glu Gln Arg Lys Val Tyr Gln Tyr Ser Asn Ser
220      420      425      430
222 Arg Ser Pro Ser Leu Glu Arg Thr Ser Lys Pro Leu Ala Pro Asn Ser
223      435      440      445
225 Asn Pro Thr Thr Lys Asp Ser Leu Leu Pro Arg Ser Pro Val Lys Pro
226      450      455      460
228 Met Asp Ser Ile Thr Ser Pro Ser Ile Ala Glu Leu His Asn Ser Val
229 465      470      475      480
231 Thr Ser Cys Asp Pro Pro Leu Arg Leu Thr Lys Ser Ser His Phe Thr
232      485      490      495
234 Asn Ile Lys Ala Val Asp Lys Leu Asp His Ser Arg Ser Asn Thr Pro
235      500      505      510
237 Ser Pro Leu Asn Leu Ser Ser Thr Ser Ser Lys Asn Ser His Ser Ser
238      515      520      525
240 Ser Tyr Thr Pro Asn Ser Phe Ser Ser Glu Glu Leu Gln Ala Glu Pro
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244 545      550      555      560
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253          595          600          605
255 Asn Asn Pro Val Phe Gly Met Asn Pro Phe Ser Ala Lys Pro Leu Tyr
256          610          615          620
258 Thr Pro Leu Pro Pro Gln Ser Ala Phe Pro Pro Ala Thr Phe Met Pro
259 625          630          635          640
261 Pro Val Gln Thr Ser Ile Pro Gly Leu Arg Pro Tyr Pro Gly Leu Asp
262          645          650          655
264 Gln Met Ser Phe Leu Pro His Met Ala Tyr Thr Tyr Pro Thr Gly Ala
265          660          665          670
267 Ala Thr Phe Ala Asp Met Gln Gln Arg Arg Lys Tyr Gln Arg Lys Gln
268          675          680          685
270 Gly Phe Gln Gly Asp Leu Leu Asp Gly Ala Gln Asp Tyr Met Ser Gly
271          690          695          700
273 Leu Asp Asp Met Thr Asp Ser Asp Ser Cys Leu Ser Arg Lys Lys Ile
274 705          710          715          720
276 Lys Lys Thr Glu Ser Gly Met Tyr Ala Cys Asp Leu Cys Asp Lys Thr
277          725          730          735
279 Phe Gln Lys Ser Ser Ser Leu Leu Arg His Lys Tyr Glu His Thr Gly
280          740          745          750
282 Lys Arg Pro His Gln Cys Gln Ile Cys Lys Lys Ala Phe Lys His Lys
283          755          760          765
285 His His Leu Ile Glu His Ser Arg Leu His Ser Gly Glu Lys Pro Tyr
286          770          775          780
288 Gln Cys Asp Lys Cys Gly Lys Arg Phe Ser His Ser Gly Ser Tyr Ser
289 785          790          795          800
291 Gln His Met Asn His Arg Tyr Ser Tyr Cys Lys Arg Glu Ala Glu Glu
292          805          810          815
294 Arg Glu Ala Ala Glu Arg Glu Ala Arg Glu Lys Gly His Leu Gly Pro
295          820          825          830
297 Thr Glu Leu Leu Met Asn Arg Ala Tyr Leu Gln Ser Ile Thr Pro Gln
298          835          840          845
300 Gly Tyr Ser Asp Ser Glu Glu Arg Glu Ser Met Pro Arg Asp Gly Glu
301          850          855          860
303 Ser Glu Lys Glu His Glu Lys Glu Gly Glu Glu Gly Tyr Gly Lys Leu
304 865          870          875          880
306 Arg Arg Arg Asp Gly Asp Glu Glu Glu Glu Glu Glu Glu Glu Ser
307          885          890          895
309 Glu Asn Lys Ser Met Asp Thr Asp Pro Glu Thr Ile Arg Asp Glu Glu
310          900          905          910
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